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MULTIDIMENSIONAL ULTRASOUND DOPPLER SIGNAL ANALYSIS FOR FETAL ACTIVITY MONITORING

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Abstract—Fetal activity parameters such as movements, heart rate and the related parameters are essential indicators of fetal wellbeing, and no device provides simultaneous access to and sufficient estimation of all of these parameters to evaluate fetal health. This work was aimed at collecting these parameters to automatically separate healthy from compromised fetuses. To achieve this goal, we first developed a multi-sensor–multi-gate Doppler system. Then we recorded multidimensional Doppler signals and estimated the fetal activity parameters via dedicated signal processing techniques. Finally, we combined these parameters into four sets of parameters (or four hyperparameters) to determine the set of parameters that is able to separate healthy from other fetuses. To validate our system, a data set consisting of two groups of fetal signals (normal and compromised) was established and provided by physicians. From the estimated parameters, an instantaneous Manning-like score, referred to as the ultrasonic score, was calculated and was used together with movements, heart rate and the associated parameters in a classification process employing the support vector machine method. We investigated the influence of the sets of parameters and evaluated the performance of the support vector machine using the computation of sensibility, specificity, percentage of support vectors and total classification error. The sensitivity of the four sets ranged from 79% to 100%. Specificity was 100% for all sets. The total classification error ranged from 0% to 20%. The percentage of support vectors ranged from 33% to 49%. Overall, the best results were obtained with the set of parameters consisting of fetal movement, short-term variability, long-term variability, deceleration and ultrasound score. The sensitivity, specificity, percentage of support vectors and total classification error of this set were respectively 100%, 100%, 35% and 0%. This indicated our ability to separate the data into two sets (normal fetuses and pathologic fetuses), and the results highlight the excellent match with the clinical classification performed by the physicians. This work indicates the feasibility of detecting compromised fetuses and also represents an interesting method of close fetal monitoring during the entire pregnancy. (E-mail: kouame@irit.fr)

Key Words: Fetal monitoring, Fetal heart rhythm, Fetal movement, Multidimensional signals, Support vector machine.

INTRODUCTION

Fetal monitoring may be required at different times during pregnancy to closely monitor certain fetal and maternal disorders ([Manning et al. 1980](#)). Existing methods, which consist of asking women to count the number of fetal movements, performing a biophysical profile or Manning's test ([Manning et al. 1980](#)) or analyzing general movements, may be either subjective or time consuming. To automatically monitor fetal activity, the most commonly used system is the cardiotocogram. This device measures fetal heart rate (FHR) and

uterine contractions ([Royal College of Obstetricians and Gynecologists \[RCOG\] 2001](#)) but does not provide any information regarding fetal movements. This may explain why the performances of major classifiers, which are based on FHR variability analysis alone (through different kind of entropies), are on the order of 80% for specificity and 80% for sensitivity ([Ferrario et al. 2006](#)). We believe that additional information may improve classification performance. There are many dedicated ultrasound systems that collect additional information such as fetal movement and pseudo-breathing. Unfortunately, these systems provide only partially automated detection of fetal movement ([Karlsonn et al. 2000a](#)) or breathing ([Karlsonn et al. 2000b](#); [Yamakoshi et al. 1996](#)).

Some preliminary attempts to overcome these limitations were made by [Kribèche et al. \(2007\)](#). The main characteristic of their system is that, because of the large number of sensors needed to cover an explored area as large as possible, a large number of Doppler signals have to be processed. The aim of our work was not to develop new signal processing methods and evaluate their performance. Instead, we sought to efficiently combine some relevant existing signal processing techniques to achieve the following goals: extract fetal activity features, such as FHR and movements (and related features), and efficiently collect these features to differentiate between normal and potentially compromised fetuses. To achieve these goals, we needed first to reduce the inherent redundancy related to the large number of acquired Doppler signals and then to separate fetal from maternal signals through a blind source separation. The detailed signal processing algorithms used to extract the fetal activity parameters are not detailed here. The reader is referred to, for example, [Rouvre et al. \(2005\)](#), [Voicu \(2011\)](#) and [Voicu et al. \(2010, 2014\)](#). However, even if the number of relevant data can be reduced, a fusion step is required to provide the physical with global information, for example, an “ultrasound score,” on which a decision can be based.

Thus, the main challenge of this study, and its incremental value with respect to the previously published article ([Kribèche et al. 2007](#)), was to develop a method that enables extraction of the relevant information necessary to construct a global indicator sufficiently pertinent to classify fetuses into normal and potentially compromised groups. To validate the feasibility, this method was successfully tested on a heterogeneous data set obtained from pregnant women and composed of FHR, movement and pseudo-breathing signals. To our knowledge, this automated classification is a new and important contribution to the search for an objective method for monitoring fetal activity, which remains an open issue ([Grivell et al. 2010](#); [Kaluzynska et al. 2011](#)).

This article is arranged as follows: We first described the device produced by an industrial collaboration and used for this study, and then briefly introduce the signal processing techniques used. The ultrasound score is then introduced and investigated using fetal activity parameters obtained from real fetal Doppler signals. Finally, results from classification of sets of fetuses are discussed.

MATERIALS AND METHODS

Materials

We constructed, in collaboration with Althais Technologies (Tours, France), a multi-sensor Doppler system (Surfoetus) that was able to monitor most parts of the fetal

body. Although a complete description of the development of this device is beyond the scope of this article, we briefly provide here a brief description. The system consists of 12 ultrasound (US) sensors, each with five gates, working at a frequency of $f_0 = 2$ MHz and an electronic US device (three electronic pulsed Doppler boards and one data acquisition board). A low-noise amplifier amplifies the received signal; a deep compensated amplifier then balances the strong attenuation of the deepest gates. After complex demodulation and filtering, the Doppler components are sampled sequentially and converted. Sampling frequency, denoted F_e , is 1 kHz. Five adjustable gates ranging from 2 to 14 cm are used to explore the depths. These 12 sensors are divided into three groups of four sensors, as illustrated in [Figure 1](#). Group A (US sensors 1–4) is used to explore the fetal thorax, investigating FHR and breathing movements; group B (sensors 5–8) the upper limbs; and group C (sensors 9–12) the lower limbs. Each sensor is a circular piezoelectric element 12 mm in diameter. The ultrasound beam was not focused. A belt was used during recording to maintain all three groups.

Protocol

In this study, 44 pregnant women at a gestational age >24 weeks were prospectively enrolled. Inclusion criteria were singleton pregnancy, absence of significant maternal complications requiring premature delivery (hypertensive disorders; renal, heart or immunologic pathology; pre-existing diabetes), willingness to have pregnancy followed or delivered locally, maternal age >18 years and health insurance affiliation. Exclusion criteria were fetal malformation, maternal significant complications, pregnancy care in another center and concomitant participation in another research protocol. This study was approved by the University’s ethics committee (Clinical Investigation Centre, Innovation

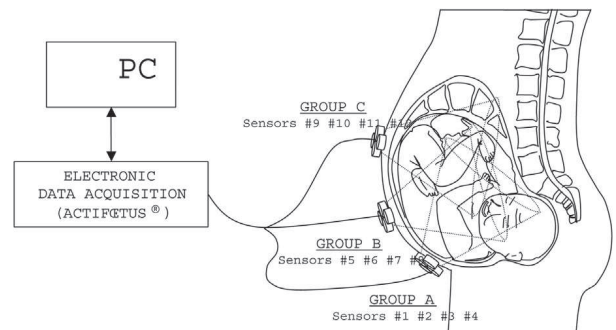


Fig. 1. Sensor group arrangement. Group A (sensors 1–4) explores the fetal thorax, investigating fetal heart rate and breathing movements. Group B (sensors 5–8) explores the upper limbs, and group C (sensors 9–12), the lower limbs. A belt is used during recording to prevent sensor movement.

Technology, Ethics). The ethics ID number was 2004-32, and the study number was CT04-SURFOETUS. All participants submitted a signed informed consent.

The standard demographic measures, expressed as the mean \pm standard deviation (range) were age = 29.9 ± 5.9 (20–44) years, height = 164 ± 6 (151–179) cm and weight = 63.23 ± 15.88 (41–123) kg. Pregnancies were classified in two groups: normal and IUGR (intra-uterine growth restriction) depending on ultrasound-estimated fetal growth at inclusion. IUGR was defined as an estimated fetal weight under the 10th percentile or a transverse abdominal perimeter below the 10th percentile for gestational age. All recordings were performed between the 24th and 40th weeks of gestation in the Obstetrics Department of Bretonneau University Hospital, Tours, France. In the “normal” group, recordings were performed every 4 weeks, and in the “IUGR” group, recordings were performed every 2 weeks. The overall mean gestational age was 30.2 weeks (median = 32 weeks). The mean gestational age of the normal group was 30.1 weeks, and the mean gestational age of the IUGR group was 29.9 weeks.

Overall, we obtained 98 Doppler signals for the 44 patients. Each examination in this protocol was 30 min long. The first group, referred to as “normal pregnancies,” consisted of 74 measurements and 31 subjects. The second group, referred to as “pathologic pregnancies” or “group with IUGR” consisted of 24 measurements and 13 subjects.

Procedures

As mentioned earlier, the electronic control system is composed of 12 sensors with five gates per sensor, and thus, 60 complex Doppler signals must be processed each millisecond. This is a very large amount of data with inevitable redundancy in the signals. Furthermore, as the signals recorded from different sensors originate from different sources of signals (fetal heart movement, maternal heart movement, respiration, *etc.*), a source separation is strongly expected to extract fetal signal only. Assuming that there is an L -dimensional zero-mean signal source vector $s(t) = [s_1(t), \dots, s_L(t)]^T$, and an M -dimensional data vector $x(t) = [x_1(t), \dots, x_M(t)]^T$ observed at each time point t . In short, L is the number of sources, and M is the number of observable mixed data. L and M are generally different. These signals can be modeled as

$$x(t) = A \cdot s(t) + u(t) \quad (1)$$

where x represents the signals observed or acquired by the sensors, referred to below as the “observation matrix”; A is the mixing matrix, corresponding to maternal body “mixing”; s is the unobservable source matrix; and $u(t)$ is a zero-mean Gaussian noise vector with a covariance

matrix Λ . In summary, we have different source signals denoted s non-observable, which are mixed and acquired by the sensors and denoted x . As stated before, the fetal signals need to be extracted and separated to correctly monitor fetal activity. To perform an unsupervised source separation, that is, without any *a priori* information, the number of independent sources must be evaluated from our data set. To do so in a parsimonious way, a redundancy information reduction must be performed.

Dimension reduction and sparse blind source separation

From the set of data, we perform dimension reduction to eliminate the data redundancy (see [Appendix A](#)). We searched for the number of independent sources using the method described in [Appendix A](#). This number of independent sources was found to be two or three. Knowing the number of sources, we performed signal separation to separate the fetal Doppler signal from the mother’s signal (see [Appendix B](#)). The last technique to be explored concerned the statement of a meta-descriptor allowing an objective evaluation of fetal state.

Ultrasound score

To compute an objective parameter including information on fetal heart rate and fetal movements, we introduced a Manning-like hyper-parameter referred to as the ultrasound score ([Ribes et al. 2011](#)). This meta-descriptor was computed every minute during each examination, as explained later. The ultrasound score is the sum of five values denoted $V1$, $V2$, $V3$, $V4$ and $V5$ that are based on the estimated fetal activity parameters, as reported in [Table 1](#).

$V1$ and $V2$ were both based on the fetal heart rate, which is an essential component in the follow-up of pregnancies ([Rouvre et al. 2005](#); [Voicu et al. 2010](#)) because it is correlated with fetal health. As reported in [Table 1](#), $V1$ represents the baseline (or mean value of FHR), and $V2$ represents the variance of the baseline. This baseline is the mean FHR computed when the variability was between 5 and 25 bpm over 1 minute. The baseline is rounded to increments of 5 bpm during a 10-min segment. The normal value lies between 110 and 160 bpm. Detection of the FHR may be improved using a specific prior ([Manning 1977](#)).

Table 1. Ultrasound score: Parameters computed each minute, associated variables and resulting points

Parameter	Variable	Points
Baseline	$V1$	1.5 to 2
Baseline variance	$V2$	–1.5 to 1
Upper limb movements	$V3$	0 to 1
Lower limb movements	$V4$	0 to 1
Offset	$V5$	2

V3 and V4 are both based on limb movements. V3 represents upper limb movements whereas V4 represents lower limb movements. These movements can be detected either on the separated magnitudes by their sudden changes or by searching for the increase or decrease in the overall phase. However, to eliminate false detection caused by phase noise, detection of fetal movement was combined with a threshold algorithm (Karlsson et al. 2000b). In this algorithm a movement was defined by a magnitude level above the noise threshold with a minimum duration of 0.2 s, followed by a minimum rest duration of 0.5 s.

Finally, V5 is an offset.

The empirical range of values was chosen on the following basis:

- If the baseline was >110 or <160 bpm, the value of V1 was set between 1.5 and 2; otherwise V1 was set at 0. More precisely, let BL denote the baseline. Thus, V1 is defined as $V1 = 0.01(BL - 110) + 1.5$ if $110 \leq BL \leq 160$, $V1 = 0$ otherwise.
- If the baseline variance was <17.5 , then the value of V2 was set between 0 and 1; if the baseline variance was >17.5 , but <22.5 , then the value of V2 was set between -1.5 and 1; otherwise, the value of V2 was set at -1.5 . More precisely, let BLV denote the baseline variance (which is a positive value). Thus V2 is defined by $V2 = BLV/17.5$ if $0 \leq BLV \leq 17.5$; $V2 = -0.5(BLV - 17.5) + 1$ if $17.5 < BLV \leq 22.5$; and $V2 = -1.5$ if $BLV > 22.5$.
- If two or more upper limb movements were detected, the value of V3 was set at 1; otherwise V3 was set at 0. If two or more lower limb movements were detected, then the value of V4 was set at 1; otherwise the value of V4 was set to 0.
- To have a positive ultrasound score, we added an offset through V5 which is set at 2.

The ultrasound score for the complete recording was the mean of the ultrasound score computed for each minute. Figure 2 illustrates an example of the ultrasound score. At the top are the mean FHR and the standard deviation for each minute. In the middle are the numbers of movements detected from the upper limbs (blue) and lower limbs (magenta). At the bottom is the ultrasound score calculated from the aforementioned parameters.

As the parameters do not have the same importance in the follow-up of fetal well-being, we investigated different sets of parameters through four new parameters referred to as hyper-parameters:

- Hyper-parameter 1 consisted of baseline, fetal movement and ultrasound score.
- Hyper-parameter 2 consisted of baseline, fetal movement, short-term variability and ultrasound score.

- Hyper-parameter 3 consisted of baseline, fetal movement and short-term variability.
- Hyper-parameter 4 consisted of fetal movement, short-term variability, long-term variability, deceleration and ultrasound score.

Note that variability is the fluctuation in the FHR, quantified as the amplitude peak-to-peak trough in beats per minute. Short-term variability (STV) of FHR is commonly used in prediction of fetal distress. STV is the oscillation of the FHR around the baseline in amplitudes of 5 to 25 bpm, measured on time windows of 3.75 s. Long-term variability is the mean value over a minute (*i.e.*, the difference between the highest and the lowest values of FHR).

Recall of used support vector machine method

To find the best hyper-parameter, we propose the use of support vector machine (SVM) classification. Different classification methods were evaluated (Duda et al. 2001), and the SVM method (Ferrario et al. 1999; Vapnik 1995) was better able to classify the current data. To ensure a better understanding, the principles of SVM classification are provided in Appendix C. In this study and using SVM principles, a score optimization problem can be reformulated as a classification problem. The problem under consideration was the study of the separability of the data (normal fetuses vs. compromised fetuses), using only estimation of the hyper-parameters mentioned earlier. We had a classification problem in which the kernel function, the set of hyper-parameters, the kernel function features and the value of the constraint C had to be chosen. The choice of kernel functions was studied empirically, and optimal results were achieved using a polynomial kernel function defined by

$$K(x, x') = (1 + x \cdot x')^d \quad (2)$$

where d is the degree defined by the user. To obtain the best results, the SVM was computed using a polynomial kernel function for different sets of hyper-parameters, kernel degrees and a constraint C defined by the user.

First, the data was pre-processed to separate the signals of the mothers from those of the fetuses, and then the parameters listed in Table 1 were computed. The effectiveness of the SVM classification was evaluated using the computation of specificity, sensitivity, percentage of support vectors and total classification accuracy defined as follows:

- Specificity is the number of true-negative decisions divided by the number of normal pregnancies.
- Sensitivity is the number of true-positive decisions divided by the number of abnormal pregnancies.

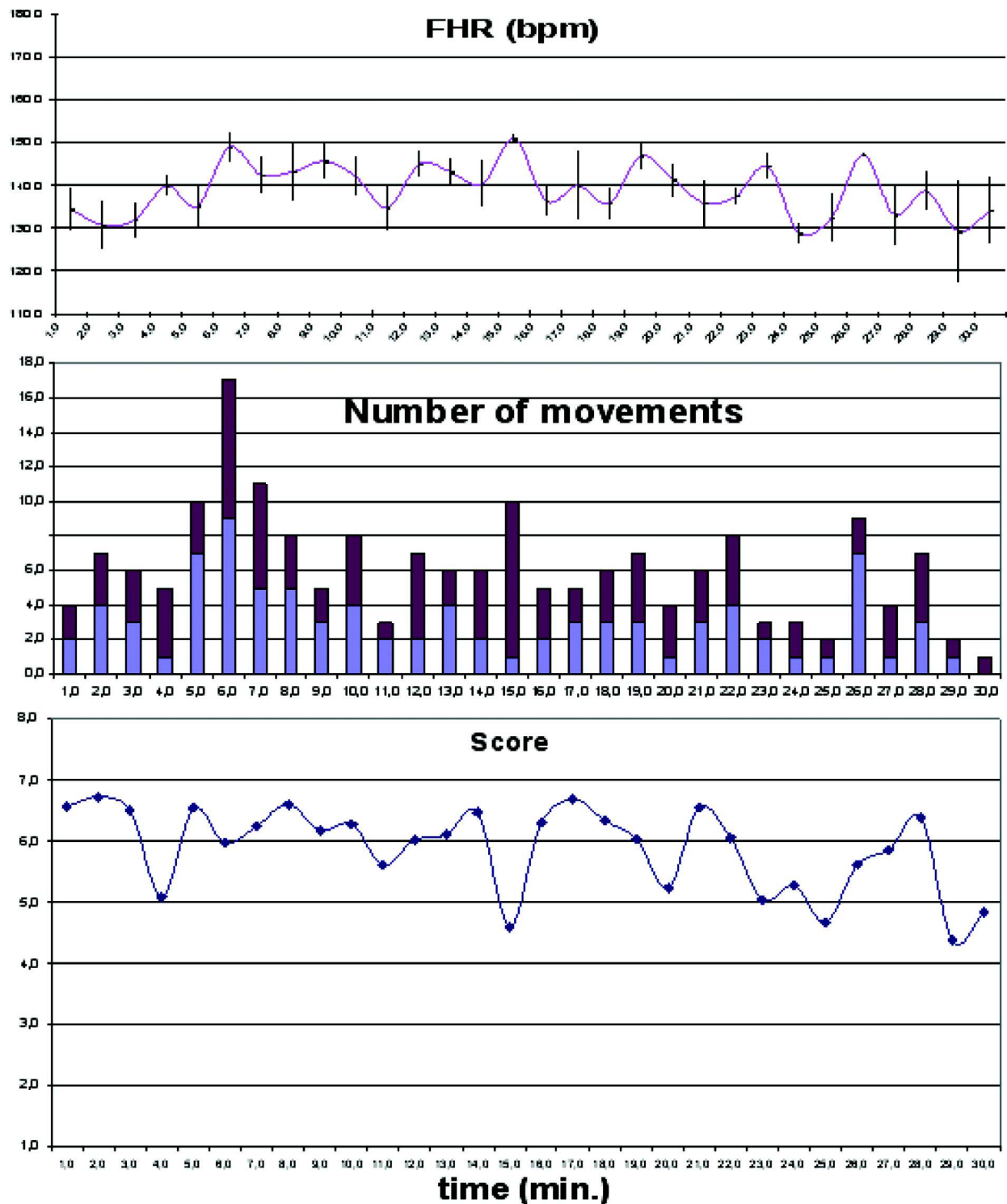


Fig. 2. Top: Mean fetal heart rate and mean STV each minute. Middle: Number of movements of the upper limbs (blue) and lower limbs (purple). Bottom: Ultrasound score calculated according to parameters. FHR = fetal heart rate; STV = short-term variability.

- Percentage of support vectors is the number of support vectors divided by the number of pregnancies.
- Classification accuracy is the number of correct decisions divided by the number of pregnancies.

A true-negative decision occurred when the classifier and the physician suggested the absence of positive detection. A true-positive decision occurred when the classifier and the physician suggested a positive

detection. As mentioned previously, a binary clinical classification (normal pregnancies and pregnancies associated with IUGR disorders) was performed by physicians.

RESULTS

We first searched for the best hyper-parameter. For different kernel degrees d and different constraints C , we performed the classification process and compared classification accuracies. Figure 3 illustrates the relationship between total error and constraint for the different hyper-parameters: for all hyper-parameters, total error decreased when constraint C increased. Hyper-parameters 2 and 3 differed only by consideration of the ultrasound score, and hyper-parameter 2, which contained this parameter, always yielded greater accuracy than hyper-parameter 3, confirming that inclusion of the ultrasound score facilitated and improved the classification process. For all constraint C values, hyper-parameter 4 always yielded the best results, and we chose to investigate only this hyper-parameter in the remainder of our study.

To find the optimal degree of the polynomial kernel function, we evaluated the performance of the classification considering hyper-parameter 4 and increasing constraint C for different degrees, as illustrated in Figure 4. Higher degrees offered more general solutions, with a reduction in the number of support vectors, projecting data into a higher-dimensional space and providing a lower error rate. At the same time, if we chose to use high levels of constraint C , we obtained better classification accuracy. The trade-off between the complexity of the decision region and the training error rate can be monitored by changing parameter C .

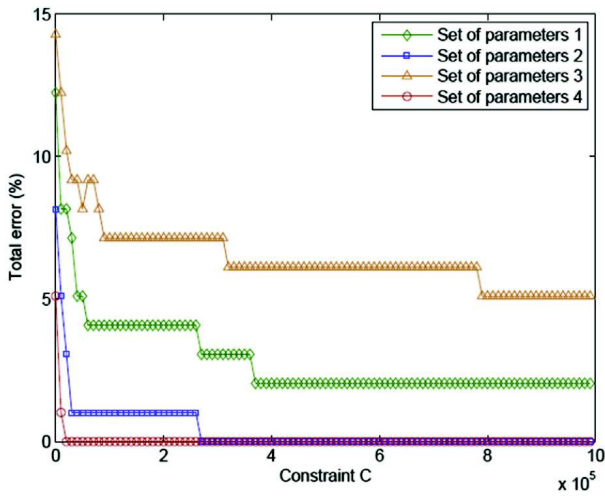


Fig. 3. Comparison of the total error for all hyper-parameters and various constraints for a degree value $d = 15$. Hyper-parameter 4 was always better.

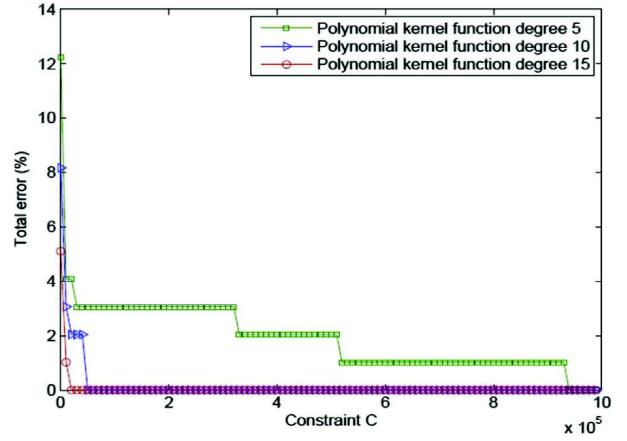


Fig. 4. For various values of constraints and various degrees of kernel, the total error made for hyper-parameter 4 is evaluated. High degrees and high constraints provided better results.

Once the effects of the degree, constraint and set of hyper-parameters had been studied, the results summarized in Tables 2 and 3 were obtained for a degree value $d = 15$ and a constraint value $C = 9.0 \times 10^5$ for all sets of hyper-parameters. Table 2 outlines the distribution of normal pregnancies, and Table 3, the distribution of pregnancies that presented with IUGR disorders. Zones 1–4, revealed by the classifier, corresponded to areas related to the hyper-planes found, as illustrated in Figure 5. Zones 1 and 4 are the areas in which the fetuses were classified without any doubt as normal and compromised, respectively. Zones 2 and 3 provided trends and were associated with probable normality for zone 2 and probable compromise for zone 3. It can be seen in Table 2, for all sets of hyper-parameters, that the normal pregnancies were found only in zones 1 and 2, that is, in the zones of normality. In Table 3, for the sets of hyper-parameters 2 and 4, it can be seen that no pregnancy was classified in zones 1 and 2; there were no false-negative decisions. From these two tables, we can

Table 2. Distribution of normal pregnancies, polynomial kernel function, degree $d = 15$, constraint $C = 9.0 \times 10^5$ *

Zone	Hyper-parameter			
	1	2	3	4
1	75.67%	68.91%	64.86%	68.91%
2	24.32%	31.08%	35.13%	31.08%
3	0	0	0	0
4	0	0	0	0

* Zones 1–4 were provided by the classifier and corresponded to a location related to the three hyperplanes. Zones 1 and 4 were the zones where the fetuses were clearly classified as normal fetuses and as fetuses presenting intrauterine growth retardation disorders, respectively. Zones 2 and 3 provided trends and were associated with probable normality for zone 2 and probable disorder for zone 3.

Table 3. Distribution of the pregnancies presenting with disorders, polynomial kernel function, degree $d = 15$, constraint $C = 9.0 \times 10^5$ *

Zone	Hyper-parameter			
	1	2	3	4
1	4.16%	0	8.33%	0
2	4.16%	0	12.5%	0
3	50%	58.33%	70.83%	50
4	41.66%	41.66%	8.33%	50

* Zones 1–4 were provided by the classifier and corresponded to location related to the three hyperplanes. Zones 1 and 4 were the zones where the fetuses were clearly classified as normal fetuses and as fetuses presenting with intrauterine growth retardation disorders, respectively. Zones 2 and 3 provided trends and were associated with probable normality for zone 2 and probable pathology for zone 3.

conclude that hyper-parameters 1–4 can identify all normal pregnancies (Table 2), and hyper-parameters 2 and 4 can identify all pregnancies that present with an IUGR disorder (Table 3).

It should be noted that we were able to identify all normal pregnancies (Table 2) and all pathologic pregnancies (Table 3), and these were good results concerning the separability of the data. Table 4 provides the overall results using evaluation of sensitivity and specificity. It can be seen that all sets of hyper-parameters, including the ultrasound score, yielded better results. Sensitivity, specificity and total error rate were excellent for hyper-parameters 2 and 4, which confirms the separability of the data in a high-dimensional space. According to the percentage of support vectors, we suggest use of hyper-parameter 4, which used fewer support vectors during the learning stage. Note also that the specificity and sensitivity values included the support vector.

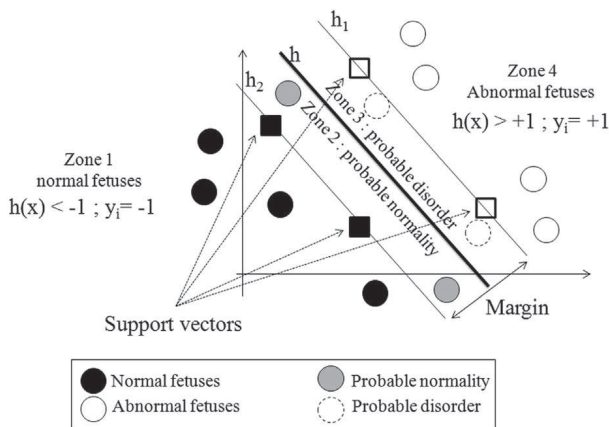


Fig. 5. Support vector machine classification. Zones 1–4 are provided by the classifier and correspond to areas related to the hyperplanes. Zones 1 and 4 are the areas where the fetuses were classified respectively as normal fetuses and as abnormal fetuses. Zones 2 and 3 provide respectively trends of probable normality and probable disorder.

Table 4. Overall evaluation: the total error, the percentage of support vectors, the sensitivity and the specificity were computed for all hyper-parameters

	Hyper-parameter			
	1	2	3	4
Total error	20.4%	0	5.1%	0
Support vectors	32.65%	37.75%	48.97%	35.71%
Sensitivity	91.66%	100%	79.16%	100%
Specificity	100%	100%	100%	100%

DISCUSSION AND CONCLUSIONS

We have illustrated in this work that combining some efficient and dedicated signal processing techniques for extraction of fetal activity features (fetal heart rhythm, variabilities, fetal limb, accelerations) can enable close separation of normal from compromised fetuses. To do so we introduced an ultrasound score, including the parameters fetal heart rate value, short-term variability and numbers of movements of the upper and lower limbs. We combined this score and other fetal activity parameters to obtain different hyper-parameters. Using these hyper-parameters and a SVM classification method, we illustrated the ability of our system to separate data into two sets, normal pregnancies and pathologic pregnancies, and obtained excellent matches to clinical classifications performed by physicians. These are valuable results and indicate an interesting way forward for home fetal monitoring. To our knowledge, this system is unique.

This study had two main limitations that will be investigated in our future works: binary classification and limited validation. With respect to binary classification, we considered here two groups (normal and IUGR). Different types of pathologies and subgroups of pathologies, with different stages, could be valuable in testing the robustness of our method. As for limited validation, the study was performed in a single clinical center. Using data from different clinical centers and performing classification of new and independent data would also help to enhance the validity of the methods on a large scale.

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APPENDIX A

DIMENSION REDUCTION

We investigated different methods of estimating the number of sources in an observation matrix such as the maximum description length, Bayesian information criterion and Akaike information criterion (AIC), which was the most suited to our data set (see also Ikeda and Toyama 2000). For convenience, model (1) can be rewritten in the framework of factor analysis as

$$x = Bf + \mu + \varepsilon \quad (\text{A.1})$$

where f is normally distributed as $f \sim N(0, I_m)$ and I_m is the $m \times m$ identity matrix; ε is normally distributed as $\varepsilon \sim N(0, \Sigma)$; and Σ is a diagonal matrix $L \times L$. f and ε are mutually independent. Signals are analyzed in short stationary windows, and therefore, because of the stationarity of ε , μ and f , for convenience in eqn (A.1) we voluntarily ignore the time variable t . The mean of x is given by μ , which is assumed here to be zero. The optimum number of sources is then estimated using the AIC.

$$(\hat{B}, \hat{\Sigma})_{MLE} = \arg\max_{B, \Sigma} L(B, \Sigma) \quad (\text{A.2})$$

where C_x is the covariance matrix of x and

$$L(B, \Sigma) = -\frac{1}{2} \left\{ \text{tr} \left(C_x (\Sigma + BB^T)^{-1} \right) + \log(\det(\Sigma + BB^T)) + L \log(2\pi) \right\}$$

For a set of data $x(t = 1, \dots, N)$, the AIC is then defined as

$$AIC = -L(\hat{B}, \hat{\Sigma}) + \frac{2}{N} \left(L(m+1) - \frac{m(m-1)}{2} \right) \quad (\text{A.3})$$

where

$$m \leq Q = \frac{1}{2} \left\{ 2L + 1 - \sqrt{8L + 1} \right\} \quad (\text{A.4})$$

To estimate the number of sources in the mixture x , we first estimated B and Σ for $1 \leq m \leq Q$, m being defined by eqn (A.4). The correct number of sources m was the one that minimized the AIC. The correct number of sources is important *a priori* information to perform source separation which gives the dimensions of the mixing matrix A to be evaluated.

Once the redundancy was reduced, the next important step was to separate the different sources in an unsupervised manner.

APPENDIX B

SPARSE BLIND SOURCE SEPARATION

Independent component analysis (ICA) is one of the best known techniques used for blind source separation. It aims at recovering unobserved sources from an available mixture observed by sensors. In other words, ICA makes it possible to separate independent data from a set of observable data. Many algorithms have been developed to perform ICA (e.g., Maxkurt [Cardoso 1999], JADE [Cardoso 1999], Infomax [Touretzky et al. 1996], fastica [Bingham and Hyvarinen 2000; Cardoso 1998; Lee 1998]).

Independent component analysis finds a linear coordinate system (the unmixing system) such that the resulting signals are statistically independent. Derived from (1) and not taking noise into account (i.e., $u(t) = 0$) is the ICA model

$$x(t) = A.s(t) \quad (\text{A.5})$$

where A is an $M \times L$ scalar mixing matrix, in which M is the number of observations and L is the number of sources recovered. The goal of ICA

is to find a linear transformation W of the observed matrix x that makes the output as independent as possible:

$$z(t) = W.x(t) = W.A.s(t) \quad (\text{A.6})$$

where z is an estimate of the sources. The sources are recovered when W is the exact inverse of A up to a permutation, sign and scale change. These methods work well when the noise level is low (typically a signal-to-noise ratio > 20 dB); their advantage is that they are the fastest techniques. However, because of the presence of high noise in our application, we used techniques including noise in the mixture model. To reduce ICA drawbacks in terms of noise, Attias (1999) proposed a model in which the source distributions are a mixture of Gaussians, the parameters of which are to be estimated jointly with the mixing matrix. Moreover, previous studies on ICA and blind source separation assume that the source distributions are sparse. Sparsity is the case where the data can be represented by a very small number of coefficients; that is, most of the source data coefficients are close to zero. Applying a transformation such as discrete cosine transform or wavelet transform to the original data can be considered sparse. Sparsity can improve ICA for two reasons: first, the statistical accuracy with which the mixing matrix can be estimated is related to how non-Gaussian the source distributions are; second, given A , the quality of the source estimates is also better for sparser sources.

Davis and Mitinoudis (2004) proposed a method that assumes the source distributions to be sparse using Mallat's modified discrete cosine transform (Mallat 1998). This technique referred to as sparse blind source separation uses two Gaussian vectors to model the distribution of source coefficients: one "on" state corresponding to a high-variance Gaussian vector, and one "off" state corresponding to a small variance. It estimates the mixing matrix, the noise covariance and the weights in the mixture of Gaussians with an expectation maximization-based procedure. Source recovery is performed using the least mean square method.

APPENDIX C

SUPPORT VECTOR MACHINE CLASSIFICATION

Given the training data set $S = \{x_i, y_i\}$, with $x_i \in \mathbf{R}^d$ the training vectors and $y_i = \{-1; +1\}$ the associated classes, we used the canonical form for SVM, which searches for the optimal hyperplane characterized by a normal w and an offset b that gives the maximum margin and satisfies the constraint

$$y_i(w \cdot x_i + b) \geq 1 - \xi_i \quad i = 1, 2, \dots \quad (\text{A.7})$$

where the ξ_i are a set of positive slack variables, and " \cdot " is the inner product. Each vector may be at a distance $\xi_i / \|w\|$ on the wrong side of the margin hyperplane. The ξ_i are a measure of the misclassification error, and the penalty function is then given by

$$C \sum_{i=1}^n \xi_i \quad (\text{A.8})$$

where C is a parameter chosen by the user. A larger C corresponds to a higher penalty for errors. It is a convex optimization problem, the aim of which is to find the global minimum. Through reformulation of the problem using the Lagrangian multiplier method, the solution is given by the saddle point of the Lagrange function

$$L_p = \frac{1}{2} \|w\|^2 + C \sum_{i=1}^n \xi_i - \sum_{i=1}^n \alpha_i (y_i(w \cdot x_i + b) - 1 + \xi_i) - \sum_{i=1}^n \beta_i \xi_i \quad (\text{A.9})$$

where the (α_i, β_i) terms are a set of non-negative Lagrangian multipliers. The minimum of the Lagrangian function is computed with respect to w and b . By setting the respective derivative and substituting these solutions, we obtain the dual objective function:

$$L_D = \frac{-1}{2} \sum_{i=1}^n \sum_{j=1}^n \alpha_i \alpha_j y_i y_j \mathbf{x}_i \cdot \mathbf{x}_j + \sum_{i=1}^n \alpha_i \quad (\text{A.10})$$

The minimum solution of the primal problem can be obtained by maximizing the dual objective function:

$$\tilde{\alpha} = \operatorname{argmax}_{\alpha} \left(\frac{-1}{2} \sum_{i=1}^n \sum_{j=1}^n \alpha_i \alpha_j y_i y_j \mathbf{x}_i \cdot \mathbf{x}_j + \sum_{i=1}^n \alpha_i \right) \quad (\text{A.11})$$

$$\text{s.t. } C \geq \alpha_i \geq 0 \quad \forall i \quad \text{and} \quad \sum_{i=1}^n \alpha_i y_i = 0 \quad (\text{A.12})$$

The support vectors are the data points denoted \mathbf{x}_s for which $y_i(\mathbf{w} \cdot \mathbf{x} + \mathbf{b}) - 1 + \xi_i = 0$ and $\alpha_i \neq 0$. They lie at a distance $\xi_i / \|\mathbf{w}\|$ on the wrong side of the margin hyperplane or on the margin hyperplanes denoted h_1, h_2 if $\xi_i = 0$. The optimal solution for the normal vector \mathbf{w} is exclusively defined by the set of support vectors, and it can be written as a linear combination of these support vectors:

$$\tilde{\mathbf{w}} = \sum_{i=1}^n \tilde{\alpha}_i y_i \mathbf{x}_i \quad (\text{A.13})$$

The optimal solution for the offset b can be obtained with

$$\tilde{b} = y_s - \sum_{i=1}^n \tilde{\alpha}_i y_i (\mathbf{x}_i \cdot \mathbf{x}_s) \quad (\text{A.14})$$

The optimal separating hyperplane is then given by

$$h(\mathbf{x}) = \operatorname{sign}(\tilde{\mathbf{w}} \cdot \mathbf{x} + \tilde{\mathbf{b}}) \quad (\text{A.15})$$

where $\operatorname{sign}(\cdot)$ is the sign function. The optimal linear boundary with the largest margin in the input space can be found using a restricted number of points called support vectors, which guarantee sufficient generalization power and robust behavior. In the case where a linear boundary in the input space is ineffective, the original space can be mapped into a high-dimensional space using a non-linear function, and the problem can be solved in this enlarged space. The mapping process is based on the chosen kernel K function satisfying Mercer's condition that

$$K(\mathbf{x}_i, \mathbf{x}_j) = \Phi(\mathbf{x}_i) \cdot \Phi(\mathbf{x}_j) \quad (\text{A.16})$$

The input vectors appear only in the form of dot products in eqn (A.11), and by use of the kernel function in eqn (A.16), the optimization problem becomes

$$\tilde{\alpha} = \operatorname{argmax}_{\alpha} \left(\frac{-1}{2} \sum_{i=1}^n \sum_{j=1}^n \alpha_i \alpha_j y_i y_j K(\mathbf{x}_i, \mathbf{x}_j) + \sum_{i=1}^n \alpha_i \right) \quad (\text{A.17})$$

such that eqn (A.12) is true. The optimal separating hyperplane is then given by

$$h(\mathbf{x}) = \operatorname{sign} \left(\sum_{i=1}^n y_i \alpha_i K(\mathbf{x}_i, \mathbf{x}_j) + \mathbf{b} \right) \quad (\text{A.18})$$

The more general case (where a linear boundary in the input space is inappropriate using a mapping function) can be solved. It can be shown that a SVM is able to exploit very high dimensional space with strong generalization guarantees derived from the max-margin property.